

1600

# CRF Errors Edited by the STIC Systems Branch

Serial Number: 09/930,0208

CRF Edit Date: 8/24/04  
Edited by: 182



\_\_\_ Corrected the SEQ ID NO. Sequence numbers edited were:

\_\_\_\_\_

\_\_\_ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

\_\_\_\_\_

\_\_\_ Deleted:   /   invalid beginning/end-of-file text ; \_\_\_ page numbers

\_\_\_ Inserted mandatory headings/numeric identifiers, specifically:

\_\_\_\_\_

\_\_\_ Moved responses to same line as heading/numeric identifier, specifically:

\_\_\_\_\_

\_\_\_ Other:

\_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_



1600

## RAW SEQUENCE LISTING

DATE: 08/24/2004

PATENT APPLICATION: US/09/930,020B

TIME: 10:35:45

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08242004\I930020B.raw

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3 <110> APPLICANT: Gish, Kurt C.
4     Mack, David H.
5     Wilson, Keith E.
7 <120> TITLE OF INVENTION: Methods of diagnosis of colorectal cancer, compositions, and
8     methods of screening for colorectal cancer modulators
10 <130> FILE REFERENCE: 05882.0168.CPUS01
12 <140> CURRENT APPLICATION NUMBER: US 09/930,020B
13 <141> CURRENT FILING DATE: 2001-08-14
15 <150> PRIOR APPLICATION NUMBER: US 09/663,733
16 <151> PRIOR FILING DATE: 2000-09-15
18 <160> NUMBER OF SEQ ID NOS: 3
20 <170> SOFTWARE: PatentIn version 3.2
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 3375
24 <212> TYPE: DNA
25 <213> ORGANISM: Homo sapiens
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32 cctggcggtg gttcctccga cctcagccgg gtccgggtcgt gccgccctct cccaggagag      180
34 acaaacaggt gtcccacgtg gcagccgcgc cccggggcgc cctcctgtga tcccgtagcg      240
36 cccctgggcc cgagccgcgc ccgggtctgt gagtagagcc gcccgggcac cgagcgctgg      300
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40 gttttcctgt ttccagagt gccccatct ctccctctcc aggaagtcca tgtaagcaaa      420
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46 cactttgcca tcacagtctg tgacggtctg gacatcagcc ccgagagggt cagagtggga      600
48 gcattccagt tcagttccac tcctcatctg gaattccct tggattcatt ttcaacccaa      660
50 caggaagtga aggcaagaat caagaggatg gttttcaaag gagggcgcac ggagacggaa      720
52 cttgctctga aataccttct gcacagaggg ttgctggag gcagaaatgc ttctgtgccc      780
54 cagatcctca tcctgtcac tgatgggaag tcccaggggg atgtggcact gccatccaag      840
56 cagctgaagg aaaggggtgt cactgtgttt gctgtggggg tcaggtttcc caggtgggag      900
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60 gaggatgcca ccaacggcct cttcagcacc ctacagcgt cgcccatctg ctccagcgcc      1020
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70 ccagaaggac tggacggcta ccagtgcctc tgcccgtggg cttttggagg ggaggctaac      1320
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74 gggggcacca ctctggacgg cttcctgcgg gccaaagtct tcgtgaagcg gtttgtgcgg      1440
76 gccgtgctga gcgaggactc tcgggcccga gtgggtgtgg ccacatacag cagggagctg      1500
78 ctggtggcgg tgctgtggg ggagtaccag gatgtgctg acctggtctg gagcctcgat      1560

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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/930,020B

DATE: 08/24/2004

TIME: 10:35:45

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08242004\I930020B.raw

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84 ctccactgagt cacactccga ggatgaggtt gcgggcccag cgcgtcacgc aaggcgcgga 1740
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92 ctgctcttca tgttggaacac ctctgcctca gtagggcccg agaattttgc tcagatgcag 1980
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128 cgttcccttg cacacaatca atgctcgcca gaatgttgtt gacacagtaa tgcccagcag 3060
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134 cttgaggggac gtttgtgact tcttggcgac tgccctttgt gtgtggaaga gacttgaaa 3240
136 ggtctcagac tgaatgtgac caattaacca gcttggttga tgatggggga ggggctgagt 3300
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140 accttgaagg tcttc 3375

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143 &lt;210&gt; SEQ ID NO: 2

144 &lt;211&gt; LENGTH: 807

145 &lt;212&gt; TYPE: PRT

146 &lt;213&gt; ORGANISM: Homo sapiens

148 &lt;400&gt; SEQUENCE: 2

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151 1 5 10 15
154 Arg Val Pro Pro Ser Leu Pro Leu Gln Glu Val His Val Ser Lys Glu
155 20 25 30
158 Thr Ile Gly Lys Ile Ser Ala Ala Ser Lys Met Met Trp Cys Ser Ala
159 35 40 45
162 Ala Val Asp Ile Met Phe Leu Leu Asp Gly Ser Asn Ser Val Gly Lys
163 50 55 60
166 Gly Ser Phe Glu Arg Ser Lys His Phe Ala Ile Thr Val Cys Asp Gly
167 65 70 75 80
170 Leu Asp Ile Ser Pro Glu Arg Val Arg Val Gly Ala Phe Gln Phe Ser
171 85 90 95
174 Ser Thr Pro His Leu Glu Phe Pro Leu Asp Ser Phe Ser Thr Gln Gln

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DATE: 08/24/2004

PATENT APPLICATION: US/09/930,020B

TIME: 10:35:45

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08242004\I930020B.raw

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175          100          105          110
178 Glu Val Lys Ala Arg Ile Lys Arg Met Val Phe Lys Gly Gly Arg Thr
179          115          120          125
182 Glu Thr Glu Leu Ala Leu Lys Tyr Leu Leu His Arg Gly Leu Pro Gly
183          130          135          140
186 Gly Arg Asn Ala Ser Val Pro Gln Ile Leu Ile Ile Val Thr Asp Gly
187 145          150          155          160
190 Lys Ser Gln Gly Asp Val Ala Leu Pro Ser Lys Gln Leu Lys Glu Arg
191          165          170          175
194 Gly Val Thr Val Phe Ala Val Gly Val Arg Phe Pro Arg Trp Glu Glu
195          180          185          190
198 Leu His Ala Leu Ala Ser Glu Pro Arg Gly Gln His Val Leu Leu Ala
199          195          200          205
202 Glu Gln Val Glu Asp Ala Thr Asn Gly Leu Phe Ser Thr Leu Ser Ser
203          210          215          220
206 Ser Ala Ile Cys Ser Ser Ala Thr Pro Asp Cys Arg Val Glu Ala His
207 225          230          235          240
210 Pro Cys Glu His Arg Thr Leu Glu Met Val Arg Glu Phe Ala Gly Asn
211          245          250          255
214 Ala Pro Cys Trp Arg Gly Ser Arg Arg Thr Leu Ala Val Leu Ala Ala
215          260          265          270
218 His Cys Pro Phe Tyr Ser Trp Lys Arg Val Phe Leu Thr His Pro Ala
219          275          280          285
222 Thr Cys Tyr Arg Thr Thr Cys Pro Gly Pro Cys Asp Ser Gln Pro Cys
223          290          295          300
226 Gln Asn Gly Gly Thr Cys Val Pro Glu Gly Leu Asp Gly Tyr Gln Cys
227 305          310          315          320
230 Leu Cys Pro Leu Ala Phe Gly Gly Glu Ala Asn Cys Ala Leu Lys Leu
231          325          330          335
234 Ser Leu Glu Cys Arg Val Asp Leu Leu Phe Leu Leu Asp Ser Ser Ala
235          340          345          350
238 Gly Thr Thr Leu Asp Gly Phe Leu Arg Ala Lys Val Phe Val Lys Arg
239          355          360          365
242 Phe Val Arg Ala Val Leu Ser Glu Asp Ser Arg Ala Arg Val Gly Val
243          370          375          380
246 Ala Thr Tyr Ser Arg Glu Leu Leu Val Ala Val Pro Val Gly Glu Tyr
247 385          390          395          400
250 Gln Asp Val Pro Asp Leu Val Trp Ser Leu Asp Gly Ile Pro Phe Arg
251          405          410          415
254 Gly Gly Pro Thr Leu Thr Gly Ser Ala Leu Arg Gln Ala Ala Glu Arg
255          420          425          430
258 Gly Phe Gly Ser Ala Thr Arg Thr Gly Gln Asp Arg Pro Arg Arg Val
259          435          440          445
262 Val Val Leu Leu Thr Glu Ser His Ser Glu Asp Glu Val Ala Gly Pro
263          450          455          460
266 Ala Arg His Ala Arg Ala Arg Glu Leu Leu Leu Leu Gly Val Gly Ser
267 465          470          475          480
270 Glu Ala Val Arg Ala Glu Leu Glu Glu Ile Thr Gly Ser Pro Lys His
271          485          490          495

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DATE: 08/24/2004

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TIME: 10:35:45

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08242004\I930020B.raw

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274 Val Met Val Tyr Ser Asp Pro Gln Asp Leu Phe Asn Gln Ile Pro Glu
275           500           505           510
278 Leu Gln Gly Lys Leu Cys Ser Arg Gln Arg Pro Gly Cys Arg Thr Gln
279           515           520           525
282 Ala Leu Asp Leu Val Phe Met Leu Asp Thr Ser Ala Ser Val Gly Pro
283           530           535           540
286 Glu Asn Phe Ala Gln Met Gln Ser Phe Val Arg Ser Cys Ala Leu Gln
287 545           550           555           560
290 Phe Glu Val Asn Pro Asp Val Thr Gln Val Gly Leu Val Val Tyr Gly
291           565           570           575
294 Ser Gln Val Gln Thr Ala Phe Gly Leu Asp Thr Lys Pro Thr Arg Ala
295           580           585           590
298 Ala Met Leu Arg Ala Ile Ser Gln Ala Pro Tyr Leu Gly Gly Val Gly
299           595           600           605
302 Ser Ala Gly Thr Ala Leu Leu His Ile Tyr Asp Lys Val Met Thr Val
303           610           615           620
306 Gln Arg Gly Ala Arg Pro Gly Val Pro Lys Ala Val Val Val Leu Thr
307 625           630           635           640
310 Gly Gly Arg Gly Ala Glu Asp Ala Ala Val Pro Ala Gln Lys Leu Arg
311           645           650           655
314 Asn Asn Gly Ile Ser Val Leu Val Val Gly Val Gly Pro Val Leu Ser
315           660           665           670
318 Glu Gly Leu Arg Arg Leu Ala Gly Pro Arg Asp Ser Leu Ile His Val
319           675           680           685
322 Ala Ala Tyr Ala Asp Leu Arg Tyr His Gln Asp Val Leu Ile Glu Trp
323           690           695           700
326 Leu Cys Gly Glu Ala Lys Gln Pro Val Asn Leu Cys Lys Pro Ser Pro
327 705           710           715           720
330 Cys Met Asn Glu Gly Ser Cys Val Leu Gln Asn Gly Ser Tyr Arg Cys
331           725           730           735
334 Lys Cys Arg Asp Gly Trp Glu Gly Pro His Cys Glu Asn Arg Glu Trp
335           740           745           750
338 Ser Ser Cys Ser Val Cys Val Ser Gln Gly Trp Ile Leu Glu Thr Pro
339           755           760           765
342 Leu Arg His Met Ala Pro Val Gln Glu Gly Ser Ser Arg Thr Pro Pro
343           770           775           780
346 Ser Asn Tyr Arg Glu Gly Leu Gly Thr Glu Met Val Pro Thr Phe Trp
347 785           790           795           800
350 Asn Val Cys Ala Pro Gly Pro
351           805
354 <210> SEQ ID NO: 3
355 <211> LENGTH: 5
356 <212> TYPE: PRT
357 <213> ORGANISM: Homo sapiens
360 <220> FEATURE:
361 <221> NAME/KEY: misc_feature
362 <222> LOCATION: (3)..(3)
363 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
365 <400> SEQUENCE: 3

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/930,020B

DATE: 08/24/2004

TIME: 10:35:45

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08242004\I930020B.raw

W--> 367 Trp Ser Xaa Trp Ser

368 1 5

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/930,020B

DATE: 08/24/2004  
TIME: 10:35:46

Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\08242004\I930020B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 3

**VERIFICATION SUMMARY**

DATE: 08/24/2004

PATENT APPLICATION: US/09/930,020B

TIME: 10:35:46

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08242004\I930020B.raw

L:367 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0



1600

## RAW SEQUENCE LISTING

DATE: 08/23/2004

PATENT APPLICATION: US/09/930,020B

TIME: 15:58:57

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\08232004\I930020B.raw

3 <110> APPLICANT: Gish, Kurt C.  
 4 Mack, David H.  
 5 Wilson, Keith E.  
 7 <120> TITLE OF INVENTION: Methods of diagnosis of colorectal cancer, compositions, and  
 8 methods of screening for colorectal cancer modulators  
 10 <130> FILE REFERENCE: 05882.0168.CPUS01  
 12 <140> CURRENT APPLICATION NUMBER: US 09/930,020B  
 13 <141> CURRENT FILING DATE: 2001-08-14  
 15 <150> PRIOR APPLICATION NUMBER: US 09/663,733  
 16 <151> PRIOR FILING DATE: 2000-09-15  
 18 <160> NUMBER OF SEQ ID NOS: 3  
 20 <170> SOFTWARE: PatentIn version 3.2  
 22 <210> SEQ ID NO: 1  
 23 <211> LENGTH: 3375  
 24 <212> TYPE: DNA  
 25 <213> ORGANISM: Homo sapiens  
 27 <400> SEQUENCE: 1

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32	cctggcggtg	gttcctccga	cctcagccgg	gtcgggtcgt	gccgccctct	cccaggagag	180
34	acaaacaggt	gtcccacgtg	gcagccgcgc	cccggggcgc	cctcctgtga	tcccgtagcg	240
36	ccccctggcc	cgagccgcgc	ccgggtctgt	gagtagagcc	gcccgggcac	cgagcgctgg	300
38	tcgccgctct	ccttcctgta	tatcaacatg	ccccctttcc	tggtgctgga	ggccgtctgt	360
40	gttttctctg	tttccagagt	gcccccatct	ctccctctcc	aggaagtcca	tgtaagcaaa	420
42	gaaaccatcg	ggaagatttc	agctgccagc	aaaatgatgt	ggtgctcggc	tgcaaggac	480
44	atcatgtttc	tgtagatggg	gtctaacagc	gtcgggaaag	ggagctttga	aaggtccaag	540
46	cactttgcca	tcacagtctg	tgacggtctg	gacatcagcc	ccgagagggt	cagagtggga	600
48	gcattccagt	tcagttccac	tcctcatctg	gaattcccct	tggattcatt	ttcaacccaa	660
50	caggaagtga	aggcaagaat	caagaggatg	gttttcaaag	gagggcgcac	ggagacgaa	720
52	cttgctctga	aataccttct	gcacagaggg	ttgcctggag	gcagaaatgc	ttctgtgccc	780
54	cagatcctca	tcacgtctac	tgatgggaag	ttccaggggg	atgtggcact	gccatccaag	840
56	cagctgaagg	aaaggggtgt	cactgtgttt	gctgtggggg	tcaggtttcc	caggtgggag	900
58	gagctgcatg	cactggccag	cgagcctaga	gggcagcacg	tgctgttggc	tgagcaggtg	960
60	gaggatgcca	ccaacggcct	cttcagcacc	ctcagcagct	cggccatctg	ctccagcgcc	1020
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64	gagttcgctg	gcaatgcccc	atgctggaga	ggatcgcgcc	ggacccttgc	ggtgctggct	1140
66	gcacactgtc	ccttctacag	ctggaagaga	gtgttcctaa	cccaccctgc	cacctgctac	1200
68	aggaccacct	gcccaggccc	ctgtgactcg	cagccctgcc	agaatggagg	cacatgtggt	1260
70	ccagaaggac	tggacggcta	ccagtgcctc	tgcccgtggy	cctttggagg	ggaggctaac	1320
72	tgtgccctga	agctgagcct	ggaatgcagg	gtcgacctcc	tcttctgctg	ggacagctct	1380
74	gcgggcacca	ctctggacgg	cttctcgagg	gcaaaagtct	tcgtgaagcg	gtttgtgcgg	1440
76	gccgtgctga	gcgaggactc	tcgggcccga	gtgggtgtgg	ccacatacag	cagggagctg	1500
78	ctggtggcgg	tgctgtgggg	ggagtaccag	gatgtgcctg	acctggtctg	gagcctcgat	1560

Does Not Comply  
Corrected Diskette Needed

P.S

## RAW SEQUENCE LISTING

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Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\08232004\I930020B.raw

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92 ctgctcttca tgttggacac ctctgcctca gtagggcccg agaattttgc tcagatgcag 1980
94 agctttgtga gaagctgtgc cctccagttt gaggtgaacc ctgacgtgac acaggtcggc 2040
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114 tggagctctt gctctgtat tgtgagccag ggatggattc ttgagacgcc cctgaggcac 2640
116 atggctcccg tgcaggagg cagcagccgt accctccca gcaactacag agaaggcctg 2700
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120 ttcccgccgt ggcaggacc actattctca ctgaggagg aggatgtccc aactgcagcc 2820
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124 ttgatgtgta agtaataacc cactttctgt acctgctgtg ccttggttag gctatgtcat 2940
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128 cgttcctttg cacacaatca atgctcgcca gaatgttgtt gacacagtaa tgcccagcag 3060
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136 ggtctcagac tgaatgtgac caattaacca gcttggttga tgatggggga ggggctgagt 3300
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140 acctgaagg tcttc 3375

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143 &lt;210&gt; SEQ ID NO: 2

144 &lt;211&gt; LENGTH: 807

145 &lt;212&gt; TYPE: PRT

146 &lt;213&gt; ORGANISM: Homo sapiens

148 &lt;400&gt; SEQUENCE: 2

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150 Met Pro Pro Phe Leu Leu Leu Glu Ala Val Cys Val Phe Leu Phe Ser
151 1 5 10 15
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155 20 25 30
158 Thr Ile Gly Lys Ile Ser Ala Ala Ser Lys Met Met Trp Cys Ser Ala
159 35 40 45
162 Ala Val Asp Ile Met Phe Leu Leu Asp Gly Ser Asn Ser Val Gly Lys
163 50 55 60
166 Gly Ser Phe Glu Arg Ser Lys His Phe Ala Ile Thr Val Cys Asp Gly
167 65 70 75 80
170 Leu Asp Ile Ser Pro Glu Arg Val Arg Val Gly Ala Phe Gln Phe Ser
171 85 90 95
174 Ser Thr Pro His Leu Glu Phe Pro Leu Asp Ser Phe Ser Thr Gln Gln

```

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TIME: 15:58:57

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\08232004\I930020B.raw

```

175          100          105          110
178 Glu Val Lys Ala Arg Ile Lys Arg Met Val Phe Lys Gly Gly Arg Thr
179          115          120          125
182 Glu Thr Glu Leu Ala Leu Lys Tyr Leu Leu His Arg Gly Leu Pro Gly
183          130          135          140
186 Gly Arg Asn Ala Ser Val Pro Gln Ile Leu Ile Ile Val Thr Asp Gly
187 145          150          155          160
190 Lys Ser Gln Gly Asp Val Ala Leu Pro Ser Lys Gln Leu Lys Glu Arg
191          165          170          175
194 Gly Val Thr Val Phe Ala Val Gly Val Arg Phe Pro Arg Trp Glu Glu
195          180          185          190
198 Leu His Ala Leu Ala Ser Glu Pro Arg Gly Gln His Val Leu Leu Ala
199          195          200          205
202 Glu Gln Val Glu Asp Ala Thr Asn Gly Leu Phe Ser Thr Leu Ser Ser
203          210          215          220
206 Ser Ala Ile Cys Ser Ser Ala Thr Pro Asp Cys Arg Val Glu Ala His
207 225          230          235          240
210 Pro Cys Glu His Arg Thr Leu Glu Met Val Arg Glu Phe Ala Gly Asn
211          245          250          255
214 Ala Pro Cys Trp Arg Gly Ser Arg Arg Thr Leu Ala Val Leu Ala Ala
215          260          265          270
218 His Cys Pro Phe Tyr Ser Trp Lys Arg Val Phe Leu Thr His Pro Ala
219          275          280          285
222 Thr Cys Tyr Arg Thr Thr Cys Pro Gly Pro Cys Asp Ser Gln Pro Cys
223          290          295          300
226 Gln Asn Gly Gly Thr Cys Val Pro Glu Gly Leu Asp Gly Tyr Gln Cys
227 305          310          315          320
230 Leu Cys Pro Leu Ala Phe Gly Gly Glu Ala Asn Cys Ala Leu Lys Leu
231          325          330          335
234 Ser Leu Glu Cys Arg Val Asp Leu Leu Phe Leu Leu Asp Ser Ser Ala
235          340          345          350
238 Gly Thr Thr Leu Asp Gly Phe Leu Arg Ala Lys Val Phe Val Lys Arg
239          355          360          365
242 Phe Val Arg Ala Val Leu Ser Glu Asp Ser Arg Ala Arg Val Gly Val
243          370          375          380
246 Ala Thr Tyr Ser Arg Glu Leu Leu Val Ala Val Pro Val Gly Glu Tyr
247 385          390          395          400
250 Gln Asp Val Pro Asp Leu Val Trp Ser Leu Asp Gly Ile Pro Phe Arg
251          405          410          415
254 Gly Gly Pro Thr Leu Thr Gly Ser Ala Leu Arg Gln Ala Ala Glu Arg
255          420          425          430
258 Gly Phe Gly Ser Ala Thr Arg Thr Gly Gln Asp Arg Pro Arg Arg Val
259          435          440          445
262 Val Val Leu Leu Thr Glu Ser His Ser Glu Asp Glu Val Ala Gly Pro
263          450          455          460
266 Ala Arg His Ala Arg Ala Arg Glu Leu Leu Leu Leu Gly Val Gly Ser
267 465          470          475          480
270 Glu Ala Val Arg Ala Glu Leu Glu Glu Ile Thr Gly Ser Pro Lys His
271          485          490          495

```

## RAW SEQUENCE LISTING

DATE: 08/23/2004

PATENT APPLICATION: US/09/930,020B

TIME: 15:58:57

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\08232004\I930020B.raw

```

274 Val Met Val Tyr Ser Asp Pro Gln Asp Leu Phe Asn Gln Ile Pro Glu
275          500          505          510
278 Leu Gln Gly Lys Leu Cys Ser Arg Gln Arg Pro Gly Cys Arg Thr Gln
279          515          520          525
282 Ala Leu Asp Leu Val Phe Met Leu Asp Thr Ser Ala Ser Val Gly Pro
283          530          535          540
286 Glu Asn Phe Ala Gln Met Gln Ser Phe Val Arg Ser Cys Ala Leu Gln
287 545          550          555          560
290 Phe Glu Val Asn Pro Asp Val Thr Gln Val Gly Leu Val Val Tyr Gly
291          565          570          575
294 Ser Gln Val Gln Thr Ala Phe Gly Leu Asp Thr Lys Pro Thr Arg Ala
295          580          585          590
298 Ala Met Leu Arg Ala Ile Ser Gln Ala Pro Tyr Leu Gly Gly Val Gly
299          595          600          605
302 Ser Ala Gly Thr Ala Leu Leu His Ile Tyr Asp Lys Val Met Thr Val
303          610          615          620
306 Gln Arg Gly Ala Arg Pro Gly Val Pro Lys Ala Val Val Val Leu Thr
307 625          630          635          640
310 Gly Gly Arg Gly Ala Glu Asp Ala Ala Val Pro Ala Gln Lys Leu Arg
311          645          650          655
314 Asn Asn Gly Ile Ser Val Leu Val Val Gly Val Gly Pro Val Leu Ser
315          660          665          670
318 Glu Gly Leu Arg Arg Leu Ala Gly Pro Arg Asp Ser Leu Ile His Val
319          675          680          685
322 Ala Ala Tyr Ala Asp Leu Arg Tyr His Gln Asp Val Leu Ile Glu Trp
323          690          695          700
326 Leu Cys Gly Glu Ala Lys Gln Pro Val Asn Leu Cys Lys Pro Ser Pro
327 705          710          715          720
330 Cys Met Asn Glu Gly Ser Cys Val Leu Gln Asn Gly Ser Tyr Arg Cys
331          725          730          735
334 Lys Cys Arg Asp Gly Trp Glu Gly Pro His Cys Glu Asn Arg Glu Trp
335          740          745          750
338 Ser Ser Cys Ser Val Cys Val Ser Gln Gly Trp Ile Leu Glu Thr Pro
339          755          760          765
342 Leu Arg His Met Ala Pro Val Gln Glu Gly Ser Ser Arg Thr Pro Pro
343          770          775          780
346 Ser Asn Tyr Arg Glu Gly Leu Gly Thr Glu Met Val Pro Thr Phe Trp
347 785          790          795          800
350 Asn Val Cys Ala Pro Gly Pro
351          805
354 <210> SEQ ID NO: 3
355 <211> LENGTH: 5
356 <212> TYPE: PRT
357 <213> ORGANISM: Homo sapiens
360 <220> FEATURE:
361 <221> NAME/KEY: misc_feature
362 <222> LOCATION: (3)..(3)
363 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
365 <400> SEQUENCE: 3

```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/930,020B

DATE: 08/23/2004

TIME: 15:58:57

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\08232004\I930020B.raw

W--> 367 Trp Ser Xaa Trp Ser

368 1 5

377 DM\_US\8051138.v1

381 DM\_US\8051138.v1

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/930,020B

DATE: 08/23/2004  
TIME: 15:58:58

Input Set : A:\PTO.FG.txt  
Output Set: N:\CRF4\08232004\I930020B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 3

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/930,020B

DATE: 08/23/2004

TIME: 15:58:58

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\08232004\I930020B.raw

L:367 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0